

FIG 1A. PRODUCTION OF GP88 BY
TUMORIGENIC AND NON-TUMORIGENIC
CELLS

Cells

PC 3A 1246 3T3

Cell Lysate

CM

FIG 1B. GP88 mRNA EXPRESSION

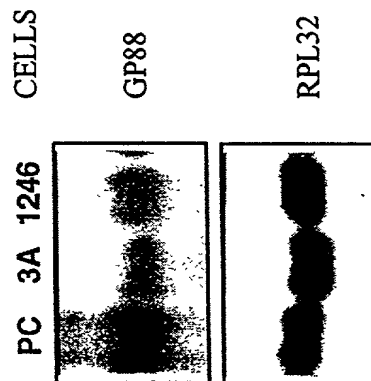
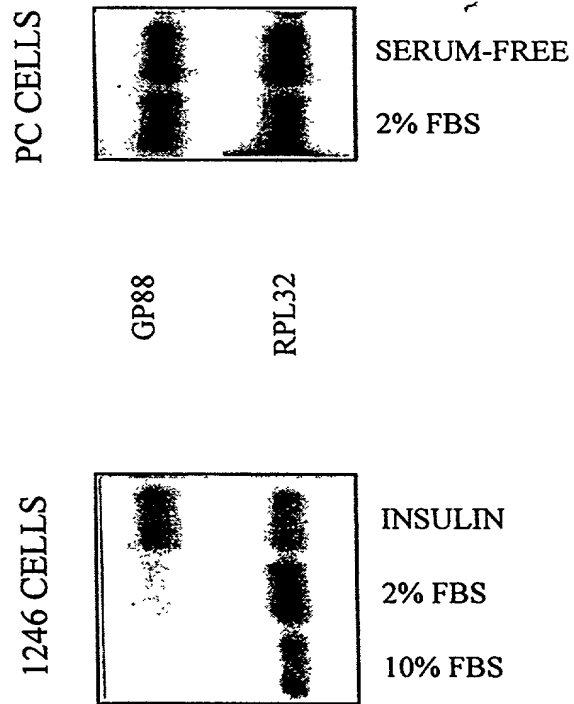


FIG 1C. GP88 mRNA EXPRESSION IN
VARIOUS CULTURE CONDITIONS



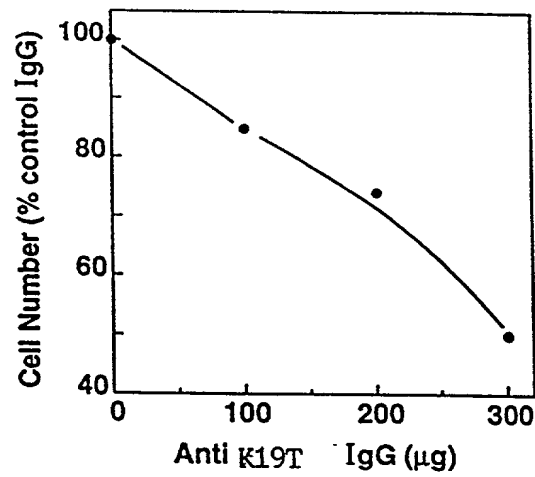
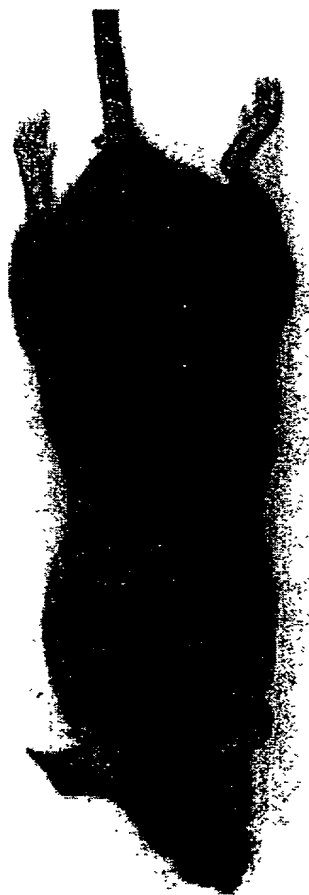


FIG. 2

FIG 3. ABSENCE OF TUMOR FORMATION IN C3H MICE BY INHIBITION OF GP88
EXPRESSION



GP88 ANTISENSE TRANSFECTED PC CELLS



CONTROL TRANSFECTED PC CELLS

FIG 4. GP88 PROTEIN EXPRESSION IN TUMOR
AND SURROUNDING TISSUES

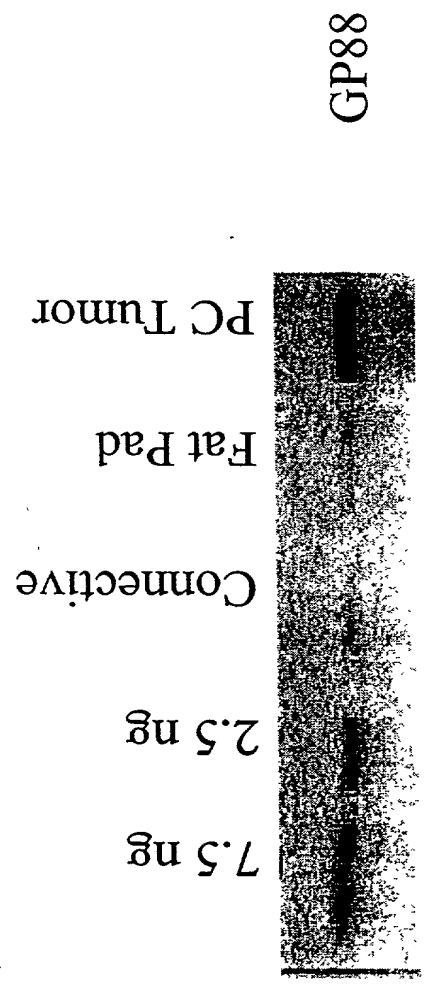
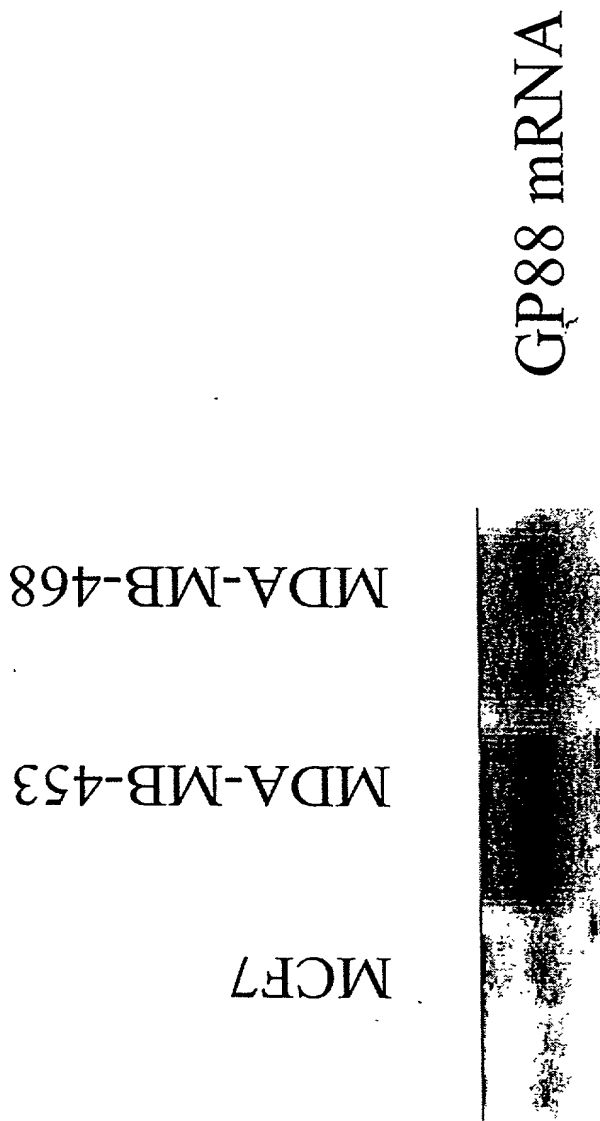


FIG 5. GP88 mRNA EXPRESSION IN
ESTROGEN-DEPENDENT AND INDEPENDENT
HUMAN MAMMARY CARCINOMA CELLS



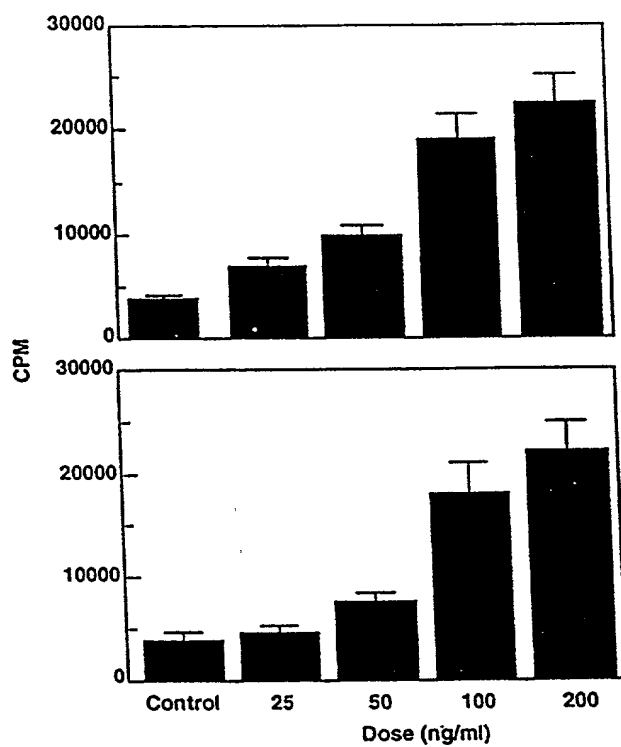
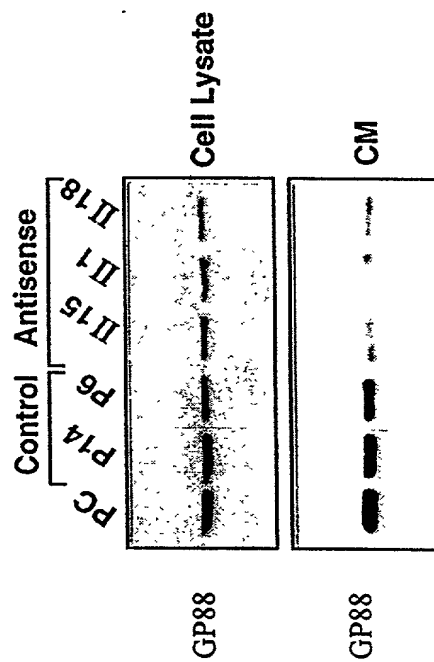


FIG. 6

FIG 7. EXPRESSION OF GP88 IN ANTISENSE AND
CONTROL TRANSFECTED PC CELLS



Mouse GP88 cDNA

[GGA	CCC	CGA	CGC	AGA	CAG	ACC	ATG	TGG	GTC	CTG	ATG	AGC	TGG	CTG	46
								M	W	V	L	M	S	W	L	8
GCC	TTC	GCG	GCA	GGG	CTG	GTA	GCC	GGA	ACA	CAG	TGT	CCA	GAT	GGG	CAG	94
A	F	A	A	G	L	V	A	G	T	Q	C	P	D	G	Q	24
TTC	TGC	CCT	GTT	GCC	TGC	TGC	CTT	GAC	CAG	GGA	GGA	GCC	AAC	TAC	AGC	142
F	C	P	V	A	C	C	L	D	Q	G	G	A	N	Y	S	40
TGC	TGT	AAC	CCT	CTT	CTG	GAC	ACA	TGG	CCT	AGA	ATA	ACG	AGC	CAT	CAT	190
C	C	N	P	L	L	D	T	W	P	R	I	T	S	H	H	56
CTA	GAT	GGC	TCC	TGC	CAG	ACC	CAT	GGC	CAC	TGT	CCT	GCT	GGC	TAT	TCT	238
L	D	G	S	C	Q	T	H	G	H	C	P	A	G	Y	S	72
TGT	CTT	CTC	ACT	GTG	TCT	GGG	ACT	TCC	AGC	TGC	TGC	CCG	TTC	TCT	AAG	286
C	L	L	T	V	S	G	T	S	S	C	C	P	F	S	K	88
GGT	GTG	TCT	TGT	GGT	GAT	GGC	TAC	CAC	TGC	TGC	CCC	CAG	GGC	TTC	CAC	334
G	V	S	C	G	D	G	Y	H	C	C	P	Q	G	F	H	104
TGT	AGT	GCA	GAT	GGG	AAA	TCC	TGC	TTC	CAG	ATG	TCA	GAT	AAC	CCC	TTG	382
C	S	A	D	G	K	S	C	F	Q	M	S	D	N	P	L	120
GGT	GCT	GTC	CAG	TGT	CCT	GGG	AGC	CAG	TTT	GAA	TGT	CCT	GAC	TCT	GCC	430
G	A	V	Q	C	P	G	S	Q	F	E	C	P	D	S	A	136
ACC	TGC	TGC	ATT	ATG	GTT	GAT	GGT	TCG	TGG	GGA	TGT	TGT	CCC	ATG	CCC	478
T	C	C	I	M	V	D	G	S	W	G	C	C	P	M	P	152
CAG	GCC	TCT	TGC	TGT	GAA	GAC	AGA	GTG	CAT	TGC	TGT	CCC	CAT	GGG	GCC	526
Q	A	S	C	C	E	D	R	V	H	C	C	P	H	G	A	168
TCC	TGT	GAC	CTG	GTT	CAC	ACA	CGA	TGC	GTT	TCA	CCC	ACG	GGC	ACC	CAC	574
S	C	D	L	V	H	T	R	C	V	S	P	T	G	T	H	184
ACC	CTA	CTA	AAG	AAG	TTC	CCT	GCA	CAA	AAG	ACC	AAC	AGG	GCA	GTG	TCT	622
T	L	L	K	K	F	P	A	Q	K	T	N	R	A	V	S	200
TTG	CCT	TTT	TCT	GTC	GTG	TGC	CCT	GAT	GCT	AAG	ACC	CAG	TGT	CCC	GAT	670
L	P	F	S	V	V	C	P	D	A	K	T	Q	C	P	D	216
GAT	TCT	ACC	TGC	TGT	GAG	CTA	CCC	ACT	GGG	AAG	TAT	GGC	TGC	TGT	CCA	718
D	S	T	C	C	E	L	P	T	G	K	Y	G	C	C	P	232
ATG	CCC	AAT	GCC	ATC	TGC	TGT	TCC	GAC	CAC	CTG	CAC	TGC	TGC	CCC	CAG	766
M	P	N	A	I	C	G	S	D	H	L	H	C	C	P	Q	248
GAC	ACT	GTA	TGT	GAC	CTG	ATC	CAG	AGT	AAG	TGC	CTA	TCC	AAG	AAC	TAC	814
D	T	V	C	D	L	I	Q	S	K	C	L	S	K	N	Y	264
ACC	ACG	GAT	CTC	CTG	ACC	AAG	CTG	CCT	GGA	TAC	CCA	GTG	AAG	GAG	GTG	862
T	T	D	L	L	T	K	L	P	G	Y	P	V	K	E	V	280
AAG	TGC	GAC	ATG	GAG	GTG	AGC	TGC	CCT	GAA	GGA	TAT	ACC	TGC	TGC	CGC	910
K	C	D	M	E	V	S	C	P	E	G	Y	T	C	C	R	296
CTC	AAC	ACT	GGG	GCC	TGG	GGC	TGC	TGT	CCA	TTT	GCC	AAG	GCC	GTG	TGT	958
L	N	T	G	A	W	G	C	C	P	F	A	K	A	V	C	312

FIG. 8

Variable	Mean	SD	Min	Max
Age	34.5	10.2	18	65
Gender	Male	1.2	0	2
Marital Status	Married	1.8	0	3
Education	High School	1.5	0	3
Occupation	Unemployed	1.2	0	2
Income	\$15,000	12,000	0	40,000
Health Status	Good	1.5	0	3
Smoking	Non-smoker	1.8	0	3
Alcohol	Non-drinker	1.5	0	3
Exercise	Regular	1.2	0	2
Stress	Low	1.5	0	3
Sleep	7-8 hours	1.2	0	2
Diet	Healthy	1.8	0	3
Family Size	2-3	1.5	0	4
Neighborhood	Urban	1.2	0	2
Transportation	Car	1.5	0	3
Insurance	Health	1.8	0	3
Religion	Christian	1.5	0	3
Politics	Conservative	1.2	0	2
Interests	Sports	1.8	0	3
Travel	Domestic	1.5	0	3
Pets	Dog	1.2	0	2
Children	1-2	1.5	0	3
Parenting	Authoritative	1.8	0	3
Education	College	1.5	0	3
Occupation	Professional	1.2	0	2
Income	\$25,000	15,000	0	50,000
Health Status	Excellent	1.5	0	3
Smoking	Former	1.8	0	3
Alcohol	Occasional	1.5	0	3
Exercise	Occasional	1.2	0	2
Stress	Medium	1.5	0	3
Sleep	6-7 hours	1.2	0	2
Diet	Balanced	1.8	0	3
Family Size	3-4	1.5	0	4
Neighborhood	Suburban	1.2	0	2
Transportation	Public	1.5	0	3
Insurance	Life	1.8	0	3
Religion	Muslim	1.5	0	3
Politics	Liberal	1.2	0	2
Interests	Reading	1.8	0	3
Travel	International	1.5	0	3
Pets	Cat	1.2	0	2
Children	3-4	1.5	0	4
Parenting	Permissive	1.8	0	3
Education	Graduate	1.5	0	3
Occupation	Managerial	1.2	0	2
Income	\$35,000	20,000	0	60,000
Health Status	Very Good	1.5	0	3
Smoking	Never	1.8	0	3
Alcohol	None	1.5	0	3
Exercise	Daily	1.2	0	2
Stress	Low	1.5	0	3
Sleep	8-9 hours	1.2	0	2
Diet	Organic	1.8	0	3
Family Size	1-2	1.5	0	3
Neighborhood	Rural	1.2	0	2
Transportation	Own	1.5	0	3
Insurance	Disability	1.8	0	3
Religion	Jewish	1.5	0	3
Politics	Conservative	1.2	0	2
Interests	Gardening	1.8	0	3
Travel	None	1.5	0	3
Pets	None	1.2	0	2
Children	0	1.5	0	3
Parenting	None	1.8	0	3
Education	High School	1.5	0	3
Occupation	Unemployed	1.2	0	2
Income	\$10,000	8,000	0	20,000
Health Status	Fair	1.5	0	3
Smoking	Current	1.8	0	3
Alcohol	Regular	1.5	0	3
Exercise	None	1.2	0	2
Stress	High	1.5	0	3
Sleep	4-5 hours	1.2	0	2
Diet	Fast Food	1.8	0	3
Family Size	5-6	1.5	0	7
Neighborhood	Urban	1.2	0	2
Transportation	Public	1.5	0	3
Insurance	Medicaid	1.8	0	3
Religion	Hindu	1.5	0	3
Politics	Liberal	1.2	0	2
Interests	TV Shows	1.8	0	3
Travel	Domestic	1.5	0	3
Pets	Dog	1.2	0	2
Children	3-4	1.5	0	5
Parenting	Authoritative	1.8	0	3
Education	College	1.5	0	3
Occup				

TGT	GAG	GAT	CAC	ATT	CAT	TGC	TGC	CCG	GCA	GGG	TTT	CAG	TGT	CAC	ACA	1006
C	E	D	H	I	H	C	C	P	A	G	F	Q	C	H	T	328
GAG	AAA	GSA	ACC	TGC	GAA	ATG	GST	ATC	CTC	CAA	GTA	CCC	TGG	ATG	AAG	1054
E	K	G	T	C	E	M	G	I	L	Q	V	P	W	M	<u>K</u>	344
AAG	GTC	ATA	GCC	CCC	CTC	CGC	CTG	CCA	GAC	CCA	CAG	ATC	TTG	AAG	AGT	1102
<u>K</u>	<u>V</u>	<u>I</u>	<u>A</u>	<u>P</u>	<u>L</u>	<u>R</u>	<u>L</u>	<u>P</u>	<u>D</u>	<u>P</u>	<u>Q</u>	<u>I</u>	<u>L</u>	<u>K</u>	<u>S</u>	360
GAT	ACA	CCT	TGT	GAT	GAC	TTC	ACT	AGG	TGT	CCT	ACA	AAC	AAT	ACC	TGC	1150
<u>D</u>	<u>T</u>	<u>P</u>	<u>C</u>	<u>D</u>	<u>D</u>	<u>F</u>	<u>T</u>	<u>R</u>	<u>C</u>	<u>P</u>	<u>T</u>	<u>N</u>	<u>N</u>	<u>T</u>	<u>C</u>	376
TGC	AAA	CTC	AAT	TCT	GGG	GAC	TGG	GGC	TGC	TGT	CCC	ATC	CCA	GAG	GCT	1198
C	K	L	N	S	G	D	W	G	C	C	P	I	P	E	A	392
GTC	TGC	TGC	TCA	GAC	AAC	CAG	CAT	TGC	TGC	CCT	CAG	GGC	TTC	ACA	TGT	1246
V	C	C	S	D	N	Q	H	C	C	P	Q	G	F	T	<u>C</u>	408
CTG	GCT	CAG	GGG	TAC	TGT	CAG	AAG	GGA	GAC	ACA	ATG	GTG	GCT	GGC	CTG	1294
L	A	Q	G	Z	C	Q	K	G	D	F	M	V	A	G	L	424
GAG	AAG	ATA	CCT	GCC	CGC	CAG	ACA	ACC	CCG	CTC	CAA	ATT	GGA	GAT	ATC	1342
E	K	I	P	A	R	Q	T	T	P	L	Q	I	G	D	I	440
GGT	TGT	GAC	CAG	CAT	TGC	AGC	TGC	CCA	GTA	GGG	CAA	ACC	TGC	TGC	CCA	1390
G	C	D	<u>Q</u>	H	T	S	C	P	V	G	Q	T	C	C	P	456
AGC	CTC	AAG	GGA	AGT	TGG	GCC	TGC	TGC	CAG	CTG	CCC	CAT	GCT	GTG	TGC	1438
S	L	K	G	S	W	A	C	C	Q	L	P	H	A	V	C	472
TGT	GAG	GAC	CGG	CAG	CAC	TGT	TGC	CCG	GCC	GGG	TAC	ACC	TGC	AAC	GTG	1486
C	E	D	R	Q	H	C	C	P	A	G	Y	T	C	N	V	488
AAG	GCG	AGG	ACC	TGT	GAG	AAG	GAT	GTC	GAT	TTT	ATC	CAG	CCT	CCC	GTG	1534
K	A	R	T	C	E	K	D	V	D	F	I	Q	P	P	V	504
CTC	CTG	ACC	CTC	GGC	CCT	AAG	GTT	GGG	AAT	GTG	GAG	TGT	GGA	GAA	GGG	1582
L	L	T	L	G	P	K	V	G	N	V	E	C	G	E	G	520
CAT	TTC	TGC	CAT	GAT	AAC	CAG	ACC	TGT	TGT	AAA	GAC	AGT	GCA	GGA	GTC	1630
H	F	C	H	D	N	Q	T	C	C	K	D	S	A	G	V	536
TGG	GCC	TGC	TGT	CCC	TAC	CTA	AAG	GGT	GTC	TGC	TGT	AGA	GAT	GGA	CGT	1678
W	A	C	C	P	Y	L	K	G	V	C	C	R	D	G	R	552
CAC	TGT	TGC	CCC	GGT	GGC	TTC	CAC	TGT	<u>TCA</u>	<u>GCC</u>	<u>AGG</u>	<u>GGA</u>	<u>ACC</u>	<u>AAG</u>	<u>TGT</u>	1726
H	C	C	P	G	G	F	H	C	<u>S</u>	<u>A</u>	<u>R</u>	<u>G</u>	<u>T</u>	<u>K</u>	<u>C</u>	568
TTG	CGA	AAG	AAG	ATT	CCT	CGC	TGG	GAC	ATG	TTT	TTG	AGG	GAT	CCG	GTC	1774
<u>L</u>	<u>R</u>	<u>K</u>	<u>K</u>	<u>I</u>	<u>P</u>	<u>R</u>	<u>W</u>	<u>D</u>	<u>M</u>	<u>F</u>	<u>L</u>	<u>R</u>	<u>D</u>	<u>P</u>	<u>V</u>	584
CCA	AGA	CCG	CTA	CTG	TAA	GGA	AGG	GCT	ACA	GAC	TTA	AGG	AAC	TCC	ACA	1822
P	R	P	L	L	*											589
GTC	CTG	GGA	ACC	CTG	TTC	CGA	GGG									

A: Nucleotide sequence of human granulatin/epithelin precursor (human GP88).
Human Granulin Genbank M75161\$

[cgaggcaga ccatgtggac cttgggtgagc tgggtggcct taacagcagg gctgggtggct
ggaacgcggt gccagatgg tcagtctgc cctgtggcct gctgcctgga cccggaggga
gccagctaca gctgtgccc tccccctctg gacaaatggc ccacaacact gagcaggcat
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cactgctgcc cagggggctt cactgcagt gcagacgggc gatcctgctt ccaaagatca
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tccacgtgct gtgttatggt ctagggctcc tgggggtgct gcccctgccc ccaggcttcc
tgctgtgaag acagggtgca ctgctgtccg cagggtgcct tctgcgacct ggttcacacc
cgctgcatca caccacggg caccacccc ctggcaaga agctccctgc ccagaggact
aacagggcag tggcctgtgc cagctcgggc atgtgtccg acgcacggc cgggtgcct
gatgggtcta cctgctgtga gctgccaggt gggaagtatg gctgtgccc aatgcccaac
gccactgct gctcgcagca cctgcactgc tggccccaag acactgtgtg tgacctgac
cagagtaagt gccttccaa ggagaacgct accacggacc tctcactaa gctgcctgcg
cacacagtg gcatgtgaa atgtgacatg gaggtgagct gccagatgg ctatacctgc
tgccgtctac agtcgggggc ctggggctgc tgcctttta cccaggctgt gtgctgtgag
gaccacatac actgctgtcc cgcgggggtt acgtgtgaca cgcagaaggg tacctgtgaa
caggggcccc accagggtgc ctggatggag aaggccccag ctacactcag cctgccagac
ccacaagcct tgaagagaga tgtcccctgt gataatgtca gcagctgtcc ctctccgat
acctgctgcc aactcacgtc tggggagtgg ggctgctgc caatcccaga ggctgtctgc
tgctcgacc accagcactg ctgccccag cgatacacgt gtgtagctga ggggcagtgt
cagcgaggaa gcgagatgt ggctggactg gagaagatgc ctgcccgcg cggttccta
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tgcccgacc aggggtgggag ctgggctgc tgcagttgc cccatgctgt gtgctgcgag
gatcgccagc actgtgccc ggctggctac acctgcaacg tgaaggctcg atctgcgag
aagggaagtgg tctctgcca gcctgccacc tcttgccc tagccctca cgtgggtgtg
aaggacgtgg agtgtgggga aggcacatc tgcctgata accagacctg ctgcccagag
aacgcagagg gctgggctg ctgtccctac gccagggcg tctgtgtg tgatcgccgc
cactgtctg ctgtggctt ccgtgcgca cgcaggggta ccaagtgtt gcgaggggag
gccccgcgt gggacgccc ttgaggggac ccagcctga gacagctgct gtgagggaca
gtactgaaga ctctgcagcc ctggggacc cactcggagg gtgcctctg ctacggcctc
cctagcacct cccctaacc aaattctccc tggacccat tctgagctcc ccatccat
gggaggtgg gcctaatct aaggccctc cctgtcagaa ggggggtgag gcaaaagccc
attacaagt gccatccct cccgtttca gtggaccctg tggcagggtg ctttcccta
tccacagggg tgttgtgtg ttgggtgtgc tttaataaa gttgtact tctt*

B: Amino-acid sequence of human granulatin/epithelin precursor (human GP88).

MWTLVSWVALTAGLVAGTRCPDQGFPCVACCLDPGGASYSCCRP
LLDKWPTTLRHLGGPCQVDAHCSAGHSCFTVSGTSSCCPFPEAVACGDGHHCCPRG
FHCSADGRSCFQRSGNNSVGAIQCPDSQFECPDFSTCCVMVDGSWGCPMPQASCCED
RVHCCPHGAFCDLVHTRCITPTGTHPLAKKLPAQRTNRAVALSSVMCPDARSRCPDG
STCCELPSGKYGCCPMPNATCCSDHLHCCPDQTVCDLIQSKCLSKENATTDLLTKLPA
HTVGDVKCDMEVSCPDGYTCCRLQSGAWGCCPFTQAVCCEDHIHCCPAGFTCDTQKGT
CEQGP HQVPWMEKAPAHLSLPDQALKRDVPCDNVSSCPSSDTCCQLTSGEWGCCPIP
EAVCCSDHQHCCPQRYTCVAEGQCQRGSEIVAGLEKMPARRGSLSHPRDIGCDQHTSC
PVGGTCCPSQGGSWACCOLPHAVCCEDRQHCCPAGYTCNVKARSCEKEVVSAQPATFL
ARSPHVGVKDVCEGEGHFCHDNQTCRRDNQGWACCPYAQGVCCADRRHCCPAGFRCA
RRGTKCLRREAPRWDAPLRDPALRQLL*

Mouse GP88 protein sequence

HHVLMHSLAFAAGLVAG 17

TQCPDQGF-CPVA--CCLDQG-GAHYSCCHPLLDTHPRITSHHL 57

DGSC-QTHGCPAGY-SCLLTUSGTS-SCCPFSKGUSCGDGYHCCPQGFHCSADGKSCFQMSDHL 120

GAUQPGSQFECPSATCCIHVD-G-SHGCCPHMPQASCCEDRVHCCPHGASCDLVHTACUSPTGHTLLKKFPAQKTHRAUSLPFS 204

WUCPDAKTQCPDSDTCCCLP-TGK-VGCCPHPHNACSDHLHCCPQDTUCDLIQSKCLSKNVTDLTKLPGYPUK 278

EUKC-DHEUSCEGYTCCRLN-TGR-HGCCPFKAVUCEDHHCPCPAGFOCHTEKGTCEGILQUPHKKVUIAPRALPDQILKS 360

DTPCNOGIR-EPHNJCCKLN-SGD-HGCCPIPEAVCCSDNQHCCPQGFCTLAQGY-CQKGDTHVAGLEKIPARQITPLQIG 438

DIGCDUHT-SCVUGOTCCPSLK-G-SHACCCQLPHAVCCEDRHCCPAGVTCNUKARTCEKDUDFIQPPULLTLGPKUG 513

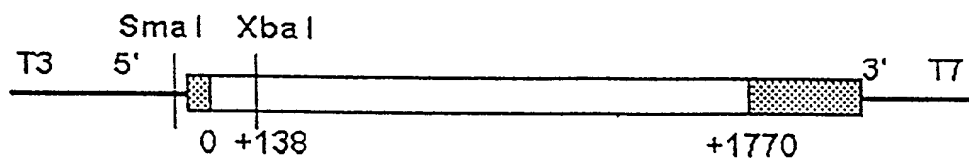
HUECGEGHF-CHDNQTCCXDSN-GU-WACCPYLKGUCCARDGRHCCPQGFHCSARGTKCLAKKIPAHDNFLRDPUPRPLL 589

consensus sequence:

C.....C.....G.....CC.....CC.....CC.D..HCCP....C.....C

1, 2: mouse epithelin 1, 2.
A, B, C, D, e, f, g: granulin A, B, C, D, E, F, G; N-terminus of granulin A, B, C, D have been sequenced.
Mouse epithelin precursor sequence is from Plowman et al.(1992).

GP88 cDNA Clone in Sk



Structure of pCMV₄ Expression Vector

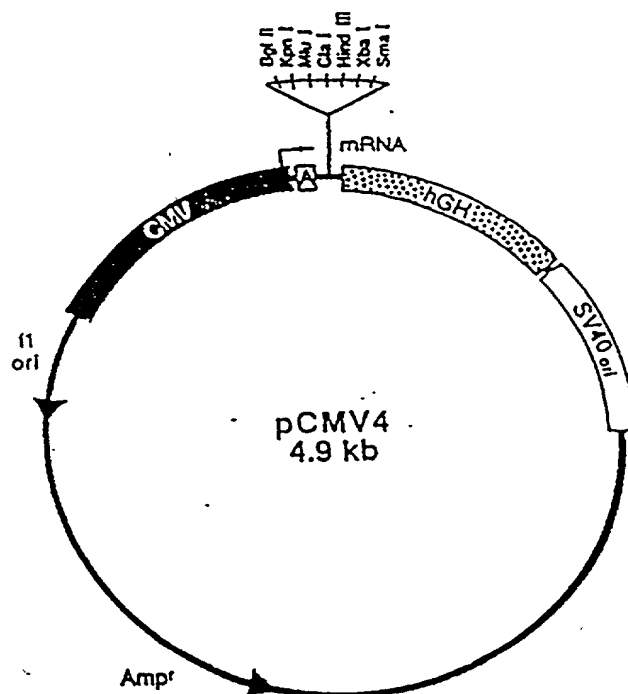


FIG. 11

FIG 12. CROSS-LINKING OF ^{125}I -rGP88 TO CCL64 CELLS

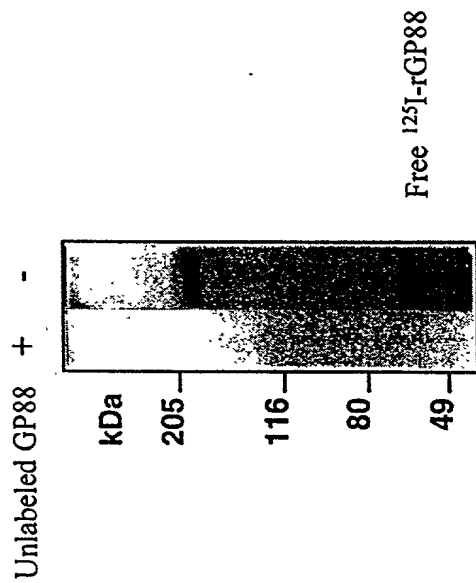
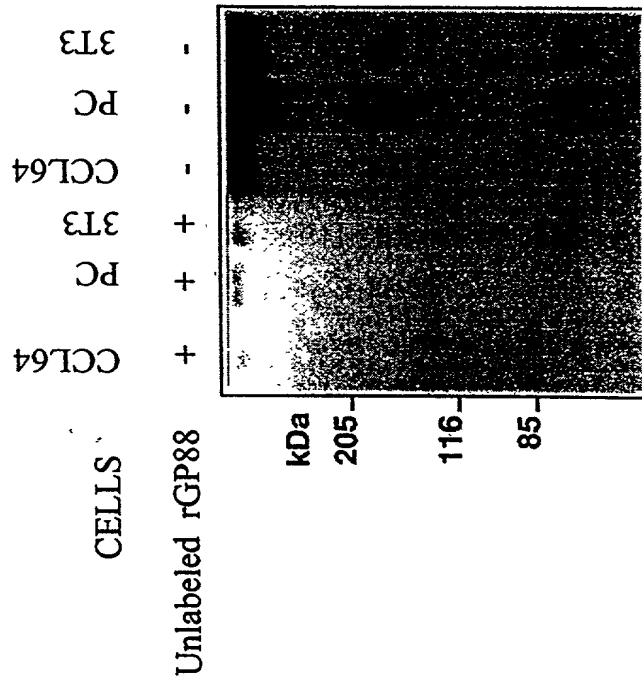
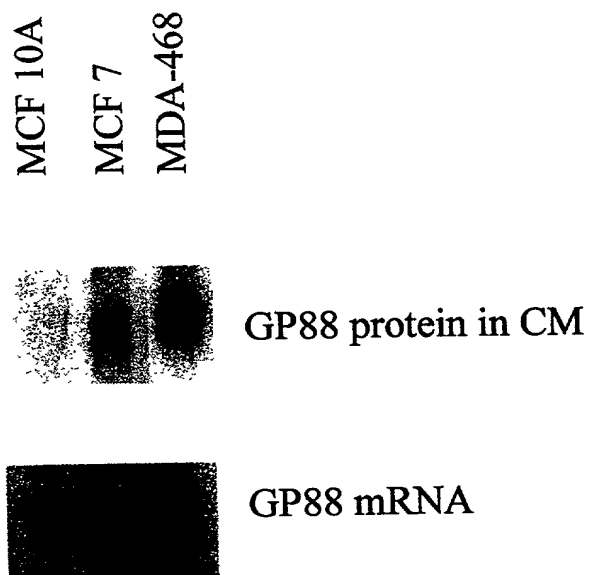


FIG 13. CROSS-LINKING OF ^{125}I -rGP88



Ad Fig 14 GP88 Expression In Non Tumorigenic (MCF 10A) And Malignant (MCF 7, MDA-468) Human Mammary Epithelial Cells



Ad Fig 15 GP88 Expression Is Inhibited By Antisense GP88 cDNA Transfection In Human Breast Carcinoma MDA-468

